IML

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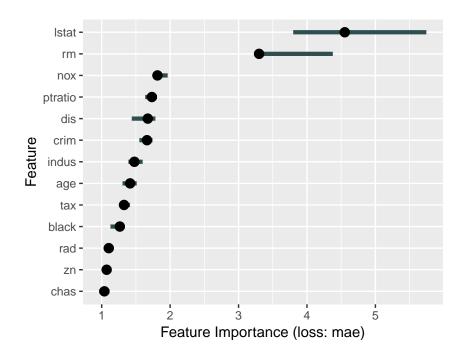
Data and Model

Fit the model on Boston housing dataset to predict the median values

```
data("Boston", package = "MASS")
head(Boston)
       crim zn indus chas
                                             dis rad tax ptratio black
                           nox
                                  rm age
## 1 0.00632 18 2.31 0 0.538 6.575 65.2 4.0900 1 296
                                                           15.3 396.90
## 2 0.02731 0 7.07
                       0 0.469 6.421 78.9 4.9671
                                                 2 242
                                                           17.8 396.90
## 3 0.02729 0 7.07 0 0.469 7.185 61.1 4.9671 2 242
                                                           17.8 392.83
## 4 0.03237 0 2.18 0 0.458 6.998 45.8 6.0622 3 222
                                                           18.7 394.63
## 5 0.06905 0 2.18 0 0.458 7.147 54.2 6.0622 3 222
                                                           18.7 396.90
## 6 0.02985 0 2.18 0 0.458 6.430 58.7 6.0622 3 222
                                                         18.7 394.12
    1stat medv
## 1 4.98 24.0
## 2 9.14 21.6
## 3 4.03 34.7
## 4 2.94 33.4
## 5 5.33 36.2
## 6 5.21 28.7
#install.packages("iml")
#library("randomForest")
library("iml")
library("randomForest")
## randomForest 4.6-12
## Type rfNews() to see new features/changes/bug fixes.
set.seed(2019)
rf = randomForest(medv ~ ., data = Boston, ntree = 1000, mtry = 4)
X = Boston[which(names(Boston) != "medv")]
predictor = Predictor$new(rf, data = X, y = Boston$medv)
```

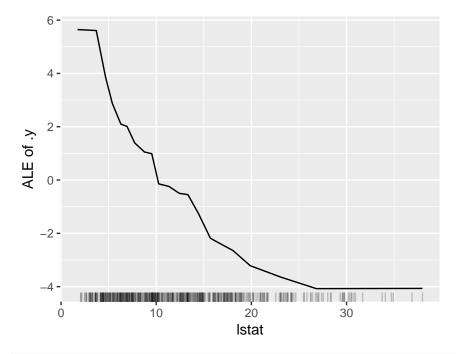
Feature importance obtained by the permutation algorithm

```
imp = FeatureImp$new(predictor, loss = "mae")
plot(imp)
```

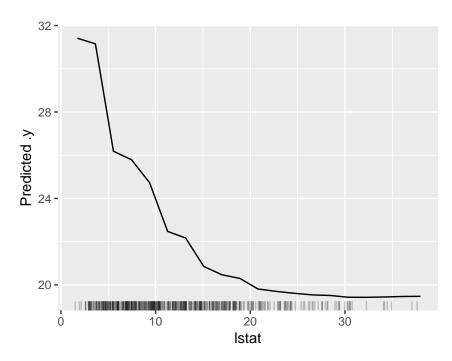


Feature effects (ALE or PDP)

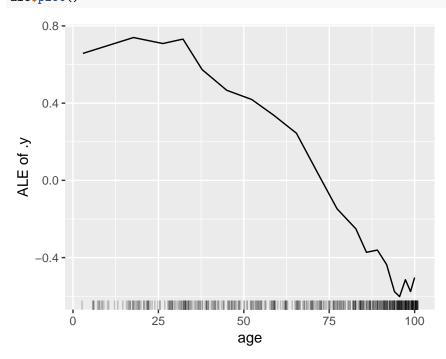
```
ale = FeatureEffect$new(predictor, feature = "lstat",method = "ale",grid.size = 20)
ale$plot()
```



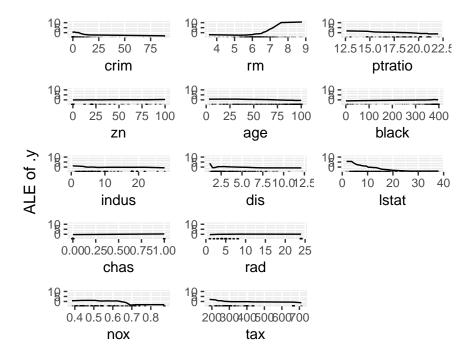
pdp = FeatureEffect\$new(predictor, feature = "lstat",method = "pdp",grid.size = 20)
pdp\$plot()



ale\$set.feature("age")
ale\$plot()

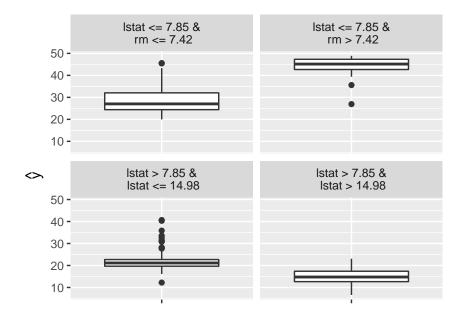


effs = FeatureEffects\$new(predictor)
plot(effs)

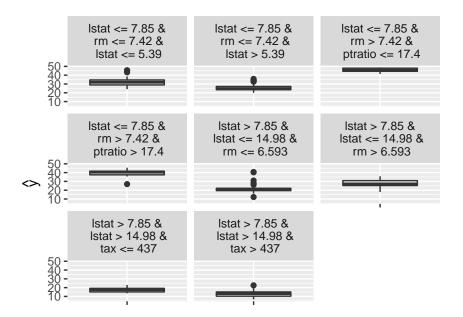


Global Surrogate Model (Tree-based)

```
tree_sur = TreeSurrogate$new(predictor,maxdepth = 2)
plot(tree_sur)
```



```
tree_sur = TreeSurrogate$new(predictor,maxdepth = 3)
plot(tree_sur)
```

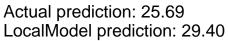


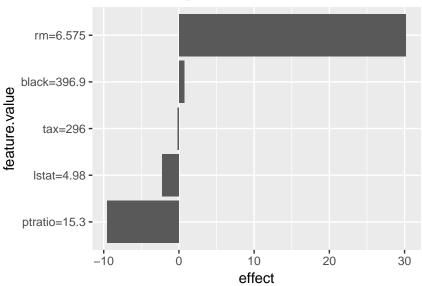
Local Surrogate Model

LocalModel fits locally weighted linear regression models to explain the indiviual predictions.

```
loc = LocalModel$new(predictor, X[1,], k = 5)
## Loading required package: glmnet
## Loading required package: Matrix
## Loading required package: foreach
## Loaded glmnet 2.0-13
## Loading required package: gower
print(loc)
## Interpretation method: LocalModel
##
##
## Analysed predictor:
## Prediction task: unknown
##
##
## Analysed data:
## Sampling from data.frame with 506 rows and 13 columns.
##
## Head of results:
##
                  beta x.recoded
                                      effect x.original feature feature.value
## rm
            4.58700691
                           6.575 30.1595704
                                                  6.575
                                                                     rm=6.575
                                                             rm
## tax
           -0.00049118
                         296.000 -0.1453893
                                                    296
                                                                      tax=296
                                                            tax
## ptratio -0.62226762
                         15.300 -9.5206946
                                                   15.3 ptratio ptratio=15.3
            0.00180294
                         396.900 0.7155868
                                                          black
## black
                                                  396.9
                                                                  black=396.9
```

```
## lstat
           -0.44052418
                            4.980 -2.1938104
                                                    4.98
                                                           lstat
                                                                     lstat=4.98
plot(loc)
```





Shapley Value

```
The contribution of a feature value to the difference between the actual prediction and the mean prediction
shapley = Shapley$new(predictor, X[1,])
print(shapley)
## Interpretation method: Shapley
## Predicted value: 25.694383, Average prediction: 22.529530 (diff = 3.164854)
##
```

Analysed predictor: ## Prediction task: unknown ##

Sampling from data.frame with 506 rows and 13 columns.

Head of results: ## feature phi.var feature.value phi ## 1 crim -0.34601096 0.898376575 crim=0.00632

rm -1.30087850 26.643823742

zn 0.02567896 0.011977925 ## 2 zn=18 ## 3 indus 0.60867790 0.584654669 indus=2.31## 4 chas -0.01097749 0.005363373 chas=0 ## 5 nox -0.27072254 0.823207605 nox=0.538## 6

plot(shapley)

Analysed data:

rm=6.575

Actual prediction: 25.69 Average prediction: 22.53

